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RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 16/533811
Source: PCT
Date Processed by STIC: 5/12/15

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PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/533,811

DATE: 05/12/2005
TIME: 08:09:33

Input Set : A:\Sequence Listing.txt
Output Set: N:\CRF4\05122005\J533811.raw

3 <110> APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA AS
 4 REPRESENTED BY THE SECRETARY OF THE DEPARTMENT OF HEALTH AND
 5 HUMAN SERVICES
 6 Valenzuela, Jesus G.
 7 Ribeiro, Jose M.C.
 8 Barral, Aldina
 9 Netto, Manoel
 10 Brodskyn, Claudia
 11 Gomes, Regis
 13 <120> TITLE OF INVENTION: LU. LONGIPALPIS POLYPEPTIDES AND METHODS OF USE
 15 <130> FILE REFERENCE: 4239-67028-08
 C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/533,811
 C--> 17 <141> CURRENT FILING DATE: 2005-04-29
 17 <150> PRIOR APPLICATION NUMBER: US 60/422,303
 18 <151> PRIOR FILING DATE: 2002-10-29
 20 <150> PRIOR APPLICATION NUMBER: PCT/US2003/034453
 21 <151> PRIOR FILING DATE: 2003-10-29
 23 <160> NUMBER OF SEQ ID NOS: 73
 25 <170> SOFTWARE: PatentIn version 3.3
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 271
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Lutzomyia longipalpis
 32 <400> SEQUENCE: 1
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 35 1 5 10 15
 38 Val Asn Ala Gln Ser Asn Tyr Cys Lys Gln Glu Ser Cys Ser Ser Gly
 39 20 25 30
 42 Gly Val Glu Arg Pro His Ile Gly Cys Lys Asn Ser Gly Asp Phe Ser
 43 35 40 45
 46 Glu Thr Cys Ser Gly Asp Ala Glu Ile Val Lys Met Asp Lys Lys
 47 50 55 60
 50 Gln Asn Leu Leu Val Lys Met His Asn Arg Leu Arg Asp Arg Phe Ala
 51 65 70 75 80
 54 Arg Gly Ala Val Pro Gly Phe Ala Pro Ala Ala Lys Met Pro Met Leu
 55 85 90 95
 58 Lys Trp Asn Asp Glu Leu Ala Lys Leu Ala Glu Tyr Asn Val Arg Thr
 59 100 105 110
 62 Cys Lys Phe Ala His Asp Lys Cys Arg Ala Ile Asp Val Cys Pro Tyr
 63 115 120 125
 66 Ala Gly Gln Asn Leu Ala Gln Met Met Ser Tyr Pro Thr His Arg Asp
 67 130 135 140
 70 Leu Asn Tyr Val Leu Lys Asn Leu Thr Arg Glu Trp Phe Trp Glu Tyr

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71	145	150	155	160
74	Arg Trp Ala Lys Gln Ser Gln Leu Asp Asn Tyr Val Gly Gly Pro Gly			
75	165	170	175	
78	Lys Asp Asn Lys Gln Ile Gly His Phe Thr Ala Phe Val His Glu Lys			
79	180	185	190	
82	Thr Asp Lys Val Gly Cys Ala Ile Ala Arg Phe Thr Asn Glu His Asn			
83	195	200	205	
86	Phe Lys Glu Thr Leu Leu Ala Cys Asn Tyr Cys Tyr Thr Asn Met Met			
87	210	215	220	
90	Lys Glu Arg Ile Tyr Thr Gln Gly Lys Pro Cys Ser Gln Cys Gln Ser			
91	225	230	235	240
94	Lys Lys Cys Gly Pro Val Tyr Lys Asn Leu Cys Asp Pro Ser Glu Lys			
95	245	250	255	
98	Val Asp Pro Thr Pro Asp Val Leu Lys Gln Trp Lys His Gly Lys			
99	260	265	270	
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103	<211> LENGTH: 905			
104	<212> TYPE: DNA			
105	<213> ORGANISM: Lutzomyia longipalpis			
107	<400> SEQUENCE: 2			
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110	tgggattgtc cgtgggtgtt aatgcacaga gcaattactg caaacaggaa tcgtgctcat			120
112	cgggaggtgt tgagagaccc catattgggt gcaaaaaactc tggagatttt tccgaaacctt			180
114	gctccggaga tgcagaaaatt gtttaagatgg acaagaagaa gcagaacctc cttgtgaaaa			240
116	tgcacaatcg cctgagagat agatttgc tgggtgcagt gccagg tttt gcaccagctg			300
118	cggaaaatgcc aatgtttaaa tggAACgtat aactggccaa attggcagag tacaacgtga			360
120	gaacgtgcaa atttgcac gataaaatgcc ggcgaattga tgtctgcccc tatgtctggac			420
122	agaatcttagc tcaaatgtat tcctatccta cccatcgaga tctaaactat gttcttaaga			480
124	atctcacaag ggaatgggtc tggagtaca gatgggctaa gcaatctcag cttgataatt			540
126	acgtgggtgg tcctggggaaa gacaacaaac aaattggaca tttcacagct tttgtgcatt			600
128	agaaaaacaga caaagtggaa tgcgctatag ctcgatttac aaatgagcac aatttttaagg			660
130	agaccctcct agcttgcac tactgctaca cgaatatgtat gaaggagagg atctacacgc			720
132	agggaaaacc ttgttcacag tgtcagagca aaaagtgtgg gccagtctac aagaacctgt			780
134	gtgatccttc ggagaagggtt gatccaactc ctgatgtcct taagcaatgg aagcatggaa			840
136	aatgattatt aagctcactt caaatgttcc caatccaaaa aaaaaaaaaa aaaaaaaaaa			900
138	aaaaaa			905
141	<210> SEQ ID NO: 3			
142	<211> LENGTH: 159			
143	<212> TYPE: PRT			
144	<213> ORGANISM: Lutzomyia longipalpis			
146	<400> SEQUENCE: 3			
148	Met Leu Leu Arg Ser Leu Phe Val Leu Phe Leu Ile Phe Leu Thr Phe			
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152	Cys Asn Ala Glu Glu Glu Leu Ile Glu Arg Lys Leu Thr Gly Lys Thr			
153	20	25	30	
156	Ile Tyr Ile Ser Thr Ile Lys Leu Pro Trp Phe Gln Ala Leu Asn His			
157	35	40	45	
160	Cys Val Lys Asn Gly Tyr Thr Met Val Ser Ile Lys Thr Phe Glu Glu			
161	50	55	60	

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Input Set : A:\Sequence Listing.txt
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164 Asn Lys Glu Leu Leu Lys Glu Leu Lys Arg Val Ile Arg Thr Glu Asp
165 65 70 75 80
168 Thr Gln Val Trp Ile Gly Gly Leu Lys His His Gln Phe Ala Asn Phe
169 85 90 95
172 Arg Trp Val Ser Asp Gly Ser His Val Ala Thr Ala Ser Gly Tyr Thr
173 100 105 110
176 Asn Trp Ala Pro Gly Glu Pro Ala Asp Ser Phe Tyr Tyr Asp Gln Phe
177 115 120 125
180 Cys Met Ala Met Leu Phe Arg Lys Asp Gly Ala Pro Trp Asp Asp Leu
181 130 135 140
184 Asn Cys Trp Val Lys Asn Leu Phe Val Cys Glu Lys Arg Asp Asp
185 145 150 155
188 <210> SEQ ID NO: 4
189 <211> LENGTH: 617
190 <212> TYPE: DNA
191 <213> ORGANISM: Lutzomyia longipalpis
193 <400> SEQUENCE: 4
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196 gcttcgttcc ttgtttgttc ttttctaata ttcttaaca ttctgcaacg ctgaggaaga 120
198 acttatttag agaaagttaa caggaaaaac gatctatac tcaacaataa agcttccgtg 180
200 gttccaagct cttaatcatt gtgttaaaaa tggctacaca atggtgtcaa ttaagacatt 240
202 tgaagagaat aaagaactcc ttaaagaact caaaaagggtg attaggacag aagatacaca 300
204 agtttggatt ggaggcctca aacatcatca atttgcaaac ttgcgttggg taagcgatgg 360
206 aagccacgta gcaacagctt cagggtacac caattgggcc ccaggggagc cagctgattc 420
208 cttctattac gatcaatttt gcatggcgat gttgttcaga aaagacggcg ctccgtggga 480
210 tgatttgaat tggtgggta agaatcttt tggttgttag aaacgagatg attgagaggc 540
212 tattttgtt atctcaccgt tttgttgaat aaaaaagaag aagaaagaca aaaaaaaaaaa 600
214 aaaaaaaaaaa aaaaaaaaaaa 617
217 <210> SEQ ID NO: 5
218 <211> LENGTH: 304
219 <212> TYPE: PRT
220 <213> ORGANISM: Lutzomyia longipalpis
222 <400> SEQUENCE: 5
224 Met Lys Leu Leu Gln Ile Ile Phe Ser Leu Phe Leu Val Phe Phe Pro
225 1 5 10 15
228 Thr Ser Asn Gly Ala Leu Thr Gly Asn Glu Ser Ala Ala Asn Ala Ala
229 20 25 30
232 Pro Leu Pro Val Val Leu Trp His Gly Met Gly Asp Ser Cys Cys Phe
233 35 40 45
236 Pro Phe Ser Leu Gly Ser Ile Lys Lys Leu Ile Glu Gln Gln Ile Pro
237 50 55 60
240 Gly Ile His Val Val Ser Leu Lys Ile Gly Lys Ser Leu Ile Glu Asp
241 65 70 75 80
244 Tyr Glu Ser Gly Phe Phe Val His Pro Asp Lys Gln Ile Gln Glu Val
245 85 90 95
248 Cys Glu Ser Leu Gln Asn Asp Leu Thr Leu Ala Asn Gly Phe Asn Ala
249 100 105 110
252 Ile Gly Phe Ser Gln Gly Ser Gln Phe Leu Arg Gly Leu Val Gln Arg
253 115 120 125

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Input Set : A:\Sequence Listing.txt
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256	Cys Ser Ser Ile Gln Val Arg Asn Leu Ile Ser Ile Gly Gly Gln His			
257	130	135	140	
260	Gln Gly Val Phe Gly Leu Pro Tyr Cys Pro Ser Leu Ser Arg Lys Thr			
261	145	150	155	160
264	Cys Glu Tyr Phe Arg Lys Leu Leu Asn Tyr Ala Ala Tyr Glu Lys Trp			
265	165	170	175	
268	Val Gln Lys Leu Leu Val Gln Ala Thr Tyr Trp His Asp Pro Leu Asn			
269	180	185	190	
272	Glu Asp Ala Tyr Arg Thr Gly Ser Thr Phe Leu Ala Asp Ile Asn Asn			
273	195	200	205	
276	Glu Arg Gln Ile Asn Asn Asp Tyr Ile Asn Asn Ile Arg Lys Leu Asn			
277	210	215	220	
280	Arg Phe Val Met Val Lys Phe Leu Asn Asp Ser Met Val Gln Pro Ile			
281	225	230	235	240
284	Glu Ser Ser Phe Phe Gly Phe Tyr Ala Pro Gly Thr Asp Thr Glu Val			
285	245	250	255	
288	Leu Pro Leu Lys Gln Ser Lys Ile Tyr Leu Glu Asp Arg Leu Gly Leu			
289	260	265	270	
292	Gln Ser Val Pro Ile Asp Tyr Leu Glu Cys Gly Gly Asp His Leu Gln			
293	275	280	285	
296	Phe Thr Lys Glu Trp Phe Ile Lys Phe Ile Ile Pro Tyr Leu Lys Gln			
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300	<210> SEQ ID NO: 6			
301	<211> LENGTH: 1273			
302	<212> TYPE: DNA			
303	<213> ORGANISM: Lutzomyia longipalpis			
305	<400> SEQUENCE: 6			
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308	ttaacaaaat tgctccattt ttgcgttttc tgaatattct gttgaaattt tgattaatct	120		
310	attttatgtg cagttttac taaaaatccc ttatcagcaa cccgggtgtct acagttttgt	180		
312	cacgctcagt agcatcttca aggtggtaag aaaaaatgaa actcctgcaa atcatcttct	240		
314	ctctcttcct ggtcttttc ccgacctcaa atggggccct gaccggaaat gaaagtgcag	300		
316	caaatgcagc tcccttgccct gtcgtcctgt ggcacgggat gggcgattct tgctgcttc	360		
318	ccttcagttt gggaaagcata aaaaaattaa ttgaacaaca aattcctggg attcatgttg	420		
320	ttagcctgaa aattggaaag tctctcattt aggactatga aagtggattt tttgttcatc	480		
322	cagacaagca aattcaggaa gtttgtgagt cacttcagaa cgatctaaca ctcgcaaatg	540		
324	gattcaatgc aattggattt tctcagggtt gtcagttcct gcgaggctt gtgcaacgat	600		
326	gttcttctat acaagtaagg aatctcattt ccattggagg acagcatcaa ggggttttg	660		
328	gtctgcccta ttgtccttcg ttgagcagaa agacttgcga atacttttaga aagctcctga	720		
330	attatgcagc ttatgaaaaa tgggtacaga aactcctagt tcaagccacc tactggcatg	780		
332	atcctctaaa tgaggatgca tatcgactg gaagcactt cttgctgat ataaataatg	840		
334	agagacaaaat caataatgac tatattaata atattcgaa gctaaatcgt tttgtgatgg	900		
336	taaagttcct caacgacagc atggttcagc caattgaatc tagttcttt ggattctacg	960		
338	ctccaggaac tgatacagaa gttctcccat taaaacaaag caagatttat ttggaagatc	1020		
340	gttgggact tcaatcagta ccgatagatt atctagaatg cgaggagat cattgcaat	1080		
342	ttacaaaaga atggttcata aagtttatca taccctatct gaagcaataa gagctgcaat	1140		
344	gtaattgatt aaaaaatgtt aaccatttca ggatgattgg gtgaccctt aaaaatataa	1200		
346	atgaaaaaat atacaaaaga aataaaattt tatattgatc ccacaaaaaa aaaaaaaaaa	1260		
348	aaaaaaaaaa aaa	1273		

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Input Set : A:\Sequence Listing.txt
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351 <210> SEQ ID NO: 7
 352 <211> LENGTH: 102
 353 <212> TYPE: PRT
 354 <213> ORGANISM: Lutzomyia longipalpis
 356 <400> SEQUENCE: 7
 358 Met Arg Asn Phe Ala Val Val Ser Leu Ala Val Ala Val Leu Leu Phe
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 362 Cys Ala Trp Pro Ile Asn Ala Glu Asp Asn Glu Glu Val Gly Lys Ala
 363 20 25 30
 366 Arg Glu Lys Arg Gly Leu Lys Asp Ala Met Glu His Phe Lys Asn Gly
 367 35 40 45
 370 Phe Lys Glu Leu Thr Lys Asp Phe Lys Leu Pro Ser Leu Pro Ser Leu
 371 50 55 60
 374 Pro Gly Phe Gly Lys Lys Pro Glu Ser Gly Ser Ser Glu Asp Ser Gly
 375 65 70 75 80
 378 Asp Lys Thr Glu Asp Thr Ser Gly Ser Lys Asp Asp Gln Ser Lys Asp
 379 85 90 95
 382 Asn Thr Val Glu Glu Ser
 383 100
 386 <210> SEQ ID NO: 8
 387 <211> LENGTH: 466
 388 <212> TYPE: DNA
 389 <213> ORGANISM: Lutzomyia longipalpis
 391 <400> SEQUENCE: 8
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 394 ctgtagtcag tttagccgtt gctgtcctgc tcttcgtgc atggcctata aatgcggaaag 120
 396 ataatgaaga agttggaaag gcgagagaaaa aaagaggctt aaaagacgca atgaaacact 180
 398 tcaaaaatgg attaaggag ctgacaaagg actttaaact tccaaagcctt ccaagtcttc 240
 400 ctggatttgg taaaaagcct gaatctggaa gttctgaaga ttctggagat aaaactgagg 300
 402 ataccagtgg atctaaggac gaccaatcaa aggataatac ggtcgaagaa tcttaagaaaa 360
 404 ggcgcaaata gctatttca aagtggcgaa tgttcttcc ttatctgaa ataaatattt 420
 406 ttaaaccttt cgaaacccaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaa 466
 409 <210> SEQ ID NO: 9
 410 <211> LENGTH: 247
 411 <212> TYPE: PRT
 412 <213> ORGANISM: Lutzomyia longipalpis
 414 <400> SEQUENCE: 9
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 417 1 5 10 15
 420 Gly Tyr Ser Trp Gln Asp Val Arg Asn Ala Asp Gln Thr Leu Trp Ala
 421 20 25 30
 424 Tyr Arg Ser Cys Gln Lys Asn Pro Glu Asp Lys Asp His Val Pro Gln
 425 35 40 45
 428 Trp Arg Lys Phe Glu Leu Pro Asp Asp Glu Lys Thr His Cys Tyr Val
 429 50 55 60
 432 Lys Cys Val Trp Thr Arg Leu Gly Ala Tyr Asn Glu Asn Glu Asn Val
 433 65 70 75 80
 436 Phe Lys Ile Asp Val Ile Thr Lys Gln Phe Asn Glu Arg Gly Leu Glu
 437 85 90 95

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 05/12/2005
PATENT APPLICATION: US/10/533,811 TIME: 08:09:34

Input Set : A:\Sequence Listing.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:50; N Pos. 1636

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/533,811

DATE: 05/12/2005

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Input Set : A:\Sequence Listing.txt

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L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:2718 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:1620